

CRF Errors Corrected by the STIC Systems Branch

Put/09
7/3/2002

Serial Number: 09/763,928A

ENTERED

CRF Processing Date:

Edited by:

Verified by:

(STIC sta)

- Changed a file from non-ASCII to ASCII
- Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- Edited a format error in the Current Application Data section, specifically:

- Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other _____.
- Added the mandatory heading and subheadings for "Current Application Data".
- Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- Changed the spelling of a mandatory field (the headings or subheadings), specifically:

- Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

- Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

- Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- Inserted colons after headings/subheadings. Headings edited included:

- Deleted extra, invalid, headings used by an applicant, specifically:

- Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file;
 page numbers throughout text; other invalid text, such as _____.
- Inserted mandatory headings, specifically: _____
- Corrected an obvious error in the response, specifically:

- Edited identifiers where upper case is used but lower case is required, or vice versa.
- Corrected an error in the Number of Sequences field, specifically:

- A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- Other:

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95



RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/763,978A

DATE: 07/03/2002
TIME: 22:40:21

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF3\07032002\I763978A.raw

P.6

3 <110> APPLICANT: Salceda, Susana
 4 Sun, Yongming
 5 Recipon, Herve
 6 Cafferkey, Robert
 8 <120> TITLE OF INVENTION: A NOVEL METHOD OF DIAGNOSING, MONITORING, STAGING, IMAGING
 9 AND TREATING VARIOUS CANCERS
 11 <130> FILE REFERENCE: DEX-0172
 13 <140> CURRENT APPLICATION NUMBER: 09/763,978A
 C--> 14 <141> CURRENT FILING DATE: 2002-04-30
 16 <150> PRIOR APPLICATION NUMBER: PCT/US99/19655
 17 <151> PRIOR FILING DATE: 1999-09-01
 19 <150> PRIOR APPLICATION NUMBER: 60/098,880
 20 <151> PRIOR FILING DATE: 1998-09-02
 22 <160> NUMBER OF SEQ ID NOS: 15
 24 <170> SOFTWARE: PatentIn version 3.1
 26 <210> SEQ ID NO: 1
 27 <211> LENGTH: 2587
 28 <212> TYPE: DNA
 29 <213> ORGANISM: Homo sapien
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 36 tggagcaatt gcactcatca ttggctttgg tatttcaggg agacactcca tcacagtcac 180
 38 tactgtcgcc tcagctggga acattgggg aatggaaatc ctgagctgca ctttgaacc 240
 40 tgacatcaa ctttctgata tcgtgataca atggctgaag gaagggttt taggcttggt 300
 42 ccatgagttc aaagaaggca aagatgagct gtcggagcag gatgaaatgt tcagaggccg 360
 44 gacagcagtg tttgctgatc aagtgatagt tggcaatgcc tctttgcggc tgaaaaacgt 420
 46 gcaactcaca gatgctggca cctacaaatg ttatatcatc acttctaaag gcaaggggaa 480
 48 tgctaacctt gagtataaaa ctggagcctt cagcatgccg gaagtgaatg tggactataa 540
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 62 gctaaaataa tgtgccttgg ccacaaaaaa gcatgcaaag tcattgttac aacagggatc 960
 64 tacagaacta tttcaccacc agatatgacc tagtttata tttctggag gaaatgaatt 1020
 66 catatctaga agtctggagt gagcaaacaa gagcaagaaaa caaaaaagaag ccaaaagcag 1080
 68 aaggctccaa tatgaacaag ataaatctat cttcaaaagac atattagaag ttggaaaaat 1140
 70 aattcatgtg aactagacaa gtgtgttaag agtgataagt aaaatgcacg tggagacaag 1200
 72 tgcattccca gatctcaggg acctccccct gcctgtcacc tggggagtgaa gaggacagga 1260
 74 tagtgcattgt tctttgtctc tgaattttta gttatatgtg ctgtaatgtt gctctgagga 1320
 76 agcccttggaa aagtctatcc caacatatcc acatcttata ttccacaaat taagctgttag 1380

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80	tagtaatggg tcaaattgatt cacttttat gatgcttcca aaggtgcctt ggcttcttt	1500
82	cccaactgac aaatgc当地 gttgagaaaa atgatcataa ttttagcata aacagagcag	1560
84	tcggcgacac cgattttata aataaactga gcacccctt tttaaacaaa caaatgc当地	1620
86	tttatttctc agatgatgtt catccgtgaa tggccaggg aaggacctt cacccgtact	1680
88	atatggcatt atgtcatcac aagctctgag gcttccttcc tccatccgtc gtggacagct	1740
90	aagacctcag tttcaatag catctagagc agtgggactc agctggggtg atttc当地	1800
92	ccatctccgg gggaaatgtct gaagacaatt ttgggttacct caatgaggg gttggaggagg	1860
94	atacagtgtct actaccaact agtggataaa ggccaggat gctgctcaac ctccattaccat	1920
96	gtacaggacg tctccccatt acaactaccc aatccgaagt gtcaactgtg tcaggactaa	1980
98	gaaaccctgg ttttagtagaaaaggccct ggaaagaggg gagccaacaa atctgtctgc	2040
100	ttctcacatt agtcatggc aaataagoat tctgtctt tggctgtgc ctccggccacag	2100
102	agagccagaa ctctatccggg caccaggata acatctctca gtgaacagag ttgacaaggc	2160
104	ctatggaaa tgcctgtatgg gattatcttc agtctgttga gcttctaagt ttcttccct	2220
106	tcattctacc ctgcaagccca agttctgtaa gagaatgcc tgagttctag ctccggcc	2280
108	cttactctga atttagatct ccagaccctt cctggccaca attcaattt aggaacacaaa	2340
110	catataccctt ccatgaagca cacacagact tttgaaagca aggacaatga ctgcttgaat	2400
112	tgaggccttg aggaatgttacttgaagga aaagaataact ttgtttccag cccccc	2460
114	acactcttca tgtgttaacc actgccttcc tggaccccttgg agccacgtg actgttattac	2520
116	atgttgttat agaaaactga ttttagagtt ctgatcgatcc aagagaatgtt taaatatac	2580
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124	<213> ORGANISM: Homo sapien	
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129	ctgcccgtca ctcggccctc ctccagccag tgctgaccag ggacttctga cctctggcc	120
131	agccaggacc tgggtgggaa ggccttcotg ctgccttggg gtgacaatct cagctccagg	180
133	ctacaggggag accgggagga tcacagagcc agcatgttac aggatccgtc cagtgatcaa	240
135	cctctgaaca gcctcgatgt caaacccctg cgccaaacccc gtatcccat ggagaccc	300
137	agaaaagggtgg ggatcccat catcatagca ctactgagcc tggccgatcatcatttg	360
139	tttgtccctca tcaaggtgtat tctggataaaa tactacttcc tctgc当地 gcctctccac	420
141	ttcatcccgaa ggaagcagct gtgtgacgga gagctggact gtc当地 ggaggacag	480
143	gagcaactgtg tcaagagctt ccccgaaaggc cctgc当地 ggactccct ctccaaaggac	540
145	cgatcccacac tgcagggtgtt ggactcggcc acagggaaact gggttctctgc ctgtttcgac	600
147	aacttcacag aagctctcgc tgagacagcc tggccgatcatc cagccaaacccc	660
149	actttcagag ctgtggagat tggcccaagac caggatctgg atgttgttga aatcacagaa	720
151	aacagccagg agcttcgtat gccaactca agtggccct gtctctcagg ctccctggc	780
153	tccctgcact gtcttgcctg tggaaagagc ctgaagaccc cccgtgtggg ggggggggag	840
155	gaggcctctg tggattcttgc ctgc当地 ggatccatcc agtacgacaa acacgtc	900
157	tgtggaggaa gcatccgttgc ccccccactgg gtc当地 gagccact gcttcaggaa	960
159	acataccgtat gtgttcaact ggaagggtgc ggc当地 gacaaactgg gcaacttccc	1020
161	atccctggct gtggccaaaga tcatcatcat tgaattcaac cccatgtacc ccaaagacaa	1080
163	tgacatcgcc ctcatgaagc tgc当地 actcaacttc tcaggccatc cagggccat	1140
165	ctgtctgccc ttctttgtat aggagctac tccagccacc ccactctggta tcattggat	1200
167	gggcttacg aagcagaatg gaggaaatgt gtctgacata ctgctgc当地 cgtcacttca	1260
169	ggtcattgtac agcacacgtt gcaatgc当地 cgtac cagggggaaag tcaccgagaa	1320
171	gatgttgc当地 gcaggcatcc cggaaaggggg tggccaggatc acagtgggtt	1380

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173	ccccctgatg taccaatctg accagtggca tgtggtggc atcgtagt gggctatgg	1440
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177	ctacaatgtc tggaaggctg agctgtaatg ctgctgcccc ttgcagtg tggagccgc	1560
179	tcccttcctg ccctgcccac ctggggatcc cccaaagtca gacacagagc aagagtcccc	1620
181	ttgggtacac ccctctgccc acagccttag catttcttgg agcagcaaag ggcctcaatt	1680
183	cctataagag accctcgag cccagaggcg cccagaggaa gtcagcagcc ctgcgtcggc	1740
185	cacacttggt gctccagca tcccaggag agacacagcc cactgaacaa ggtctcagg	1800
187	gtattgctaa gccaagaagg aactttccca cactactgaa tggaaaggcagg ctgtcttgc	1860
189	aaagcccaga tcactgtggg ctggagagga gaaggaaagg gtctgcgcca gccctgtccg	1920
191	tcttcaccca tcccaagcc tactagagca agaaaaccgt tctaataaa aatgcactgc	1980
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210	gacattgtgg ctggccacc acatggtcag cctcaaagtt gagaggctca gtaaccctcc	240
212	tatccctaga gaattccaaa gtgtggatgt aatttaacta gaaaggcatt ggtgactatc	300
214	tgtgatcctc tggaaagtatg ctatgttgc tatacttgc atccaaagcc agagggAACCC	360
216	acaatgacta gtaaaacgggt ggtctcaatg cccacttagc ctgcctct gaatttgacc	420
218	atagttggcgt tcagctgata gagcgggaag aagaaatatg catttttat gaaaaataaa	480
220	atatccaaga gaagatgaaa ctaaatggag aaattgaaat acatctactg gaagaaaaaga	540
222	tccaaattccct gaaaatgaag attgctgaga agcaaaagaca aatttgcgtg acccagaaat	600
224	tactgccagc caagagggtcc ctggatgccc acctagctgt gctccaaatt cagtttcac	660
226	agtgtacaga cagaattaaa gacctggaga aacagttcgta aacgcctgt ggtgagaata	720
228	gagctcgctt cttccaggg aaagatctga ccggaaaaaaa aatgatccaa aaatttagaca	780
230	agctggaaact acaactggcc aagaaggagg agaagctgct ggagaaggat ttcatctatg	840
232	agcaggcttc caggctcaca gacaggctct gcagcaaaac tcagggtc aagcaggaca	900
234	cactgcttt agccaagaag atgaatggct atcaaagaag gatcaaaaat gcaactgaga	960
236	aaatgatggc tcttgcgt gaggctgtcc tggaaacaagc cctaaccatt gaactccaaa	1020
238	aggaagtca gggaaaagaa gacttcatct tcacttgaa ttccaggata gaaaaaggc	1080
240	tgccactcaa taagggaaatt gagaagaat ggttggaaatg ctttcggat gaagaaatgc	1140
242	acgccttggc catcgctgaa aagtctcagg agttcttgc agcagataat cgccagctgc	1200
244	ccaaatgggt ttacacaact gcagaggcgc gtcggaaatgc ctacatccca gaagcagatg	1260
246	ccactcttcc ttggccaaaa ctttatggtg ctgggtc ttttaaaccc agtgaacctg	1320
248	gagccaatat gaggcacata aggaaacctg ttataaagcc agttgaaatc tgaatatgtg	1380
250	aacaatccca ggcctctcaa gggaaaagact tcaaccaggc ttccctgtac ccacaggtga	1440
252	aaaatgtgag cataataactt ctaatattat tgataagtaa ggttaaccaca attagtcagc	1500
254	aacagagtac aacagggttt ctatccacc accaactact ataccttca tgacgttgaa	1560
256	tgggacatag aactgtccta catttatgtc aaagttataa tttgaatcgc ttatatttc	1620
258	tttttcactt tttatattga gtacattcca gaaatttgcgt gtaggcaagg tgctataaaa	1680
260	atgcactaaa aataaatctg ttctcaatg	1709
263	<210> SEQ ID NO: 4	
264	<211> LENGTH: 257	
265	<212> TYPE: DNA	

RAW SEQUENCE LISTING

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Input Set : A:\PTO.AMC.txt

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 268 <400> SEQUENCE: 4
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 273 atatcttcta tatttaatat gaaagtctt gaaatgtatca gacagaaggg gatttcagtt 180
 275 tgcaaataat gagcaatgtt gcaattttaa cacatttcat aaatatatat tttgtcattt 240
 277 gtggagagca ccattt 257
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 281 <211> LENGTH: 359
 282 <212> TYPE: DNA
 283 <213> ORGANISM: Homo sapien
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 290 ccatcggttat cagtgcccc tggagggaca gtcacactca cttgtggctt ggctctgac 180
 292 tcagtctcta ctaatttctt ccccacctgg taccagcaga ccccaggcca ggctccacgc 240
 294 acgctcatctt acagcacaag cactcgctt tctgggtcc ctgatcgttt ctctggctcc 300
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 300 <211> LENGTH: 1372
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 302 <213> ORGANISM: Homo sapien
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 306 <222> LOCATION: (6)..(6)
 307 <223> OTHER INFORMATION: n = a, c, g or t
 310 <220> FEATURE:
 311 <221> NAME/KEY: misc_feature
 312 <222> LOCATION: (9)..(9)
 313 <223> OTHER INFORMATION: n = a, c, g or t
 316 <400> SEQUENCE: 6
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 319 ccttgggcgc cctctgtgg ccctcctgaa gctaacaggg gcgagtgtc ggtggtttac 120
 321 aaattgcctc catgcagact atgaaactgt tcagcctgtc atagtttagat ctctggcact 180
 323 ggcccaggag gtcttgcaga tttgcagatc aaggagaacc caggagttc aaagaagcgg 240
 325 ctagtaaagg tctctgagat cttgcacta gctacatcct cagggtagga ggaagatggc 300
 327 ttccagaagc atgcggctgc tcctattgtc gagctgcctg gccaacacag gagtcctgg 360
 329 tgatatcatc atgagaccca gctgtgtcc tggatggtt ttaccacaag tccaattgtc 420
 331 atggttactt caggaagctg aggaacttgtt ctgtatgcga gctcgagtgt cagtcttacg 480
 333 gaaacggagc ccacctggca tctatcctgaa gttaaagga agccagcaccc atagcagagt 540
 335 acataagtgg ctatcagaga agccagccga tatggattgg cctgcacac ccacagaaga 600
 337 ggcagcgtg gcagtggtt gatggggcca tgtatctgtc cagatcctgg tctggcaagt 660
 339 ccatgggtgg gaacaagcac tgtgtcgaga tgagctccaa taacaactt ttaacttgg 720
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 349 tctaaatgtt tgcccccgc tcccttcca cagtatcctt cttccctcctt cccctgtctc 1020
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355	ccttcagctt	ctacacc	ctt	ctgc	cctc	tccattgcct	gcaccccacc	1200
357	aactcctgct	tg	ttttcct	ttggccatgg	gaagtttac	cagtaga	atc	1260
359	tatgtgggc	catacattcc	ttaataaac	cattgtgtac	ataagaggtt	gctgtgttcc	1320	
361	agttcagtaa	atgg	gtat	tggaaaagt	aaataagacc	aagaaataca	aa	1372
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365	<211>	LENGTH:	291					
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367	<213>	ORGANISM:	Homo sapien					
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371	<222>	LOCATION:	(277)..(277)					
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378	ccactggact	tagagatg	ga	ttgaatgtt	gg	aagattaagg	aaagggagaa	120
380	gtcttaggtt	tcatcttc	ag	atgactgggt	gaacagc	agt	gttcttgc	180
382	agactaggg	aaagagcc	ag	ttctgtat	t	agcatattat	at	240
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401	<223>	OTHER INFORMATION:	n= a, c, g, or t					
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407	<223>	OTHER INFORMATION:	n= a, c, g, or t					
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413	ctatcagcta	taaaaaaaat	caactg	ccag	ccaagaactt	taaaacttta	agctgtgtat	120
415	tatagaaccg	ttttgtgt	tag	catt	ggaata	ttgtccattt	tgttaagtcat	180
417	cttaattatc	agcttgaagg	tat	ttttgt	taaaagtt	acattgaaga	acctaagtgg	240
419	atgatggat	ttggggcc	ag	tagt	gaaat	atgttcc	taaaatattt	300
421	tgg	tata	cat	gttatttta	ttat	gagatt	tgttat	360
W--> 423	tgttcc	at	ctgt	tcac	catat	gttaa	gatc	420
425	acaaaactaa	aatt	gtt	aat	taca	agaaaa	tataggtct	480
427	atacatatgg	ttgt	caca	at	cg	tata	gtat	540
429	ggtgtataaa	tttttctata	ccc	aaatt	aga	tta	ttcc	600
431	tgt	ctaattcc	tctt	cata	ttat	gtgt	ttcc	660
433	gaggacagag	gagt	ctgg	tc	tag	atgggg	aactga	720
W--> 435	aaggatgnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnn	ntaat	gtttctt	agt	780

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:6; N Pos. 6,9
Seq#:7; N Pos. 277
Seq#:8; N Pos. 410,728,729,730,731,732,733,734,735,736,737,738,739,740,741
Seq#:8; N Pos. 742,743,744,745,746,747,748,749,750,751,752,753,754,755,756
Seq#:8; N Pos. 957
Seq#:12; N Pos. 30,248,383



PCT09

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/763,978A

DATE: 06/12/2002
TIME: 14:40:15

Input Set : A:\pto.vsk.txt
Output Set: N:\CRF3\06122002\I763978A.raw

3 <110> APPLICANT: Salceda, Susana
 4 Sun, Yongming
 5 Recipon, Herve
 6 Cafferkey, Robert
 8 <120> TITLE OF INVENTION: A NOVEL METHOD OF DIAGNOSING, MONITORING, STAGING, IMAGING
 9 AND TREATING VARIOUS CANCERS
 11 <130> FILE REFERENCE: DEX-0172
 13 <140> CURRENT APPLICATION NUMBER: 09/763,978A
 C--> 14 <141> CURRENT FILING DATE: 2002-04-30
 16 <150> PRIOR APPLICATION NUMBER: PCT/US99/19655
 17 <151> PRIOR FILING DATE: 1999-09-01
 19 <150> PRIOR APPLICATION NUMBER: 60/098,880
 20 <151> PRIOR FILING DATE: 1998-09-02
 22 <160> NUMBER OF SEQ ID NOS: 15
 24 <170> SOFTWARE: PatentIn version 3.1

*Does Not Comply
Corrected Diskette Needed*

ERRORED SEQUENCES

690 <210> SEQ ID NO: 15
 691 <211> LENGTH: 492
 692 <212> TYPE: PRT
 693 <213> ORGANISM: Homo sapien
 695 <400> SEQUENCE: 15
 697 Met Ala Leu Asn Ser Gly Ser Pro Pro Ala Ile Gly Pro Tyr Tyr Glu
 698 1 5 10 15
 701 Asn His Gly Tyr Gln Pro Glu Asn Pro Tyr Pro Ala Gln Pro Thr Val
 702 20 25 30
 705 Val Pro Thr Val Tyr Glu Val His Pro Ala Gln Tyr Tyr Pro Ser Pro
 706 35 40 45
 709 Val Pro Gln Tyr Ala Pro Arg Val Leu Thr Gln Ala Ser Asn Pro Val
 710 50 55 60
 713 Val Cys Thr Gln Pro Lys Ser Pro Ser Gly Thr Val Cys Thr Ser Lys
 714 65 70 75 80
 717 Thr Lys Lys Ala Leu Cys Ile Thr Leu Thr Leu Gly Thr Phe Leu Val
 718 85 90 95
 721 Gly Ala Ala Leu Ala Ala Gly Leu Leu Trp Lys Phe Met Gly Ser Lys
 722 100 105 110
 725 Cys Ser Asn Ser Gly Ile Glu Cys Asp Ser Ser Gly Thr Cys Ile Asn
 726 115 120 125
 729 Pro Ser Asn Trp Cys Asp Gly Val Ser His Cys Pro Gly Gly Glu Asp
 730 130 135 140
 733 Glu Asn Arg Cys Val Arg Leu Tyr Gly Pro Asn Phe Ile Leu Gln Met

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734	145	150	155	160
737	Tyr Ser Ser Gln Arg Lys Ser Trp His Pro Val Cys Gln Asp Asp Trp			
738	165	170	175	
741	Asn Glu Asn Tyr Gly Arg Ala Ala Cys Arg Asp Met Gly Tyr Lys Asn			
742	180	185	190	
745	Asn Phe Tyr Ser Ser Gln Gly Ile Val Asp Asp Ser Gly Ser Thr Ser			
746	195	200	205	
749	Phe Met Lys Leu Asn Thr Ser Ala Gly Asn Val Asp Ile Tyr Lys Lys			
750	210	215	220	
753	Leu Tyr His Ser Asp Ala Cys Ser Ser Lys Ala Val Val Ser Leu Arg			
754	225	230	235	240
757	Cys Leu Ala Cys Gly Val Asn Leu Asn Ser Ser Arg Gln Ser Arg Ile			
758	245	250	255	
761	Val Gly Gly Glu Ser Ala Leu Pro Gly Ala Trp Pro Trp Gln Val Ser			
762	260	265	270	
765	Leu His Val Gln Asn Val His Val Cys Gly Gly Ser Ile Ile Thr Pro			
766	275	280	285	
769	Glu Trp Ile Val Thr Ala Ala His Cys Val Glu Lys Pro Leu Asn Asn			
770	290	295	300	
773	Pro Trp His Trp Thr Ala Phe Ala Gly Ile Leu Arg Gln Ser Phe Met			
774	305	310	315	320
777	Phe Tyr Gly Ala Gly Tyr Gln Val Gln Lys Val Ile Ser His Pro Asn			
778	325	330	335	
781	Tyr Asp Ser Lys Thr Lys Asn Asn Asp Ile Ala Leu Met Lys Leu Gln			
782	340	345	350	
785	Lys Pro Leu Thr Phe Asn Asp Leu Val Lys Pro Val Cys Leu Pro Asn			
786	355	360	365	
789	Pro Gly Met Met Leu Gln Pro Glu Gln Leu Cys Trp Ile Ser Gly Trp			
790	370	375	380	
793	Gly Ala Thr Glu Glu Lys Gly Lys Thr Ser Glu Val Leu Asn Ala Ala			
794	385	390	395	400
797	Lys Val Leu Leu Ile Glu Thr Gln Arg Cys Asn Ser Arg Tyr Val Tyr			
798	405	410	415	
801	Asp Asn Leu Ile Thr Pro Ala Met Ile Cys Ala Gly Phe Leu Gln Gly			
802	420	425	430	
805	Asn Val Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu Val Thr Ser			
806	435	440	445	
809	Asn Asn Asn Ile Trp Trp Leu Ile Gly Asp Thr Ser Trp Gly Ser Gly			
810	450	455	460	
813	Cys Ala Lys Ala Tyr Arg Pro Gly Val Tyr Gly Asn Val Met Val Phe			
814	465	470	475	480
817	Thr Asp Trp Ile Tyr Arg Gln Met Lys Ala Asn Gly			
818	485	490		

E--> 824 14

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/763,978A

DATE: 06/12/2002

TIME: 14:40:16

Input Set : A:\pto.vsk.txt

Output Set: N:\CRF3\06122002\I763978A.raw

L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:317 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:384 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:240
L:423 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:360
L:435 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:720
L:441 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:900
L:635 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:643 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:240
L:647 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:360
L:824 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:15